GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:09:48 ; Search time 1274 Seconds
                                              (without alignments)
                                        13572.847 Million cell
updates/sec

Title: US-09-300-482-311
Perfect score: 274
Sequence: 1
cttacattcctttctccacc..........gccaagcatgccgtcgaccg 274

Scoring table: IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched: 111716937 seqs, 31560637849 residues

Total number of hits satisfying chosen parameters: 223433874

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   Pending_Patents_NA_Main:*
                   1: /ABSS/Data/CRF/ptodata/1/pna/PCTUSA_COMB.seq:*
                   2: /ABSS/Data/CRF/ptodata/1/pna/PCTUSB_COMB.seq:*
                   3: /ABSS/Data/CRF/ptodata/1/pna/PCTUSC_COMB.seq:*
                   4: /ABSS/Data/CRF/ptodata/1/pna/PCTUSD_COMB.seq:*
                   5: /ABSS/Data/CRF/ptodata/1/pna/US075_COMB.seq:*
                   6: /ABSS/Data/CRF/ptodata/1/pna/US076_COMB.seq:*
                   7: /ABSS/Data/CRF/ptodata/1/pna/US077_COMB.seq:*
                   8: /ABSS/Data/CRF/ptodata/1/pna/US078_COMB.seq:*
                   9: /ABSS/Data/CRF/ptodata/1/pna/US079_COMB.seq:*
                   10: /ABSS/Data/CRF/ptodata/1/pna/US080_COMB.seq:*
                   11: /ABSS/Data/CRF/ptodata/1/pna/US081_COMB.seq:*
                   12: /ABSS/Data/CRF/ptodata/1/pna/US082_COMB.seq:*
                   13: /ABSS/Data/CRF/ptodata/1/pna/US083_COMB.seq:*
                   14: /ABSS/Data/CRF/ptodata/1/pna/US084_COMB.seq:*